CRF Error Corrected by the STIC Syst ms Branch **CRF Processing Date:** 101019.596A Serial Numb r: Edited by: Verified by: (STIC staff) Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins in cases where the sequence text was "wrapped" down to the margins are the margins and the margins are the m Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other . Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: 🗹 non-ASCII "garbage" at the beginning/end of files; 🔲 secretary initials/filename at end of file: page numbers throughout text; other invalid text, such as_____ Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected: ____ Other:

PCTIO

*Examin r: The abov corrections must be communicated to the applicant in the first Offic Action. DO NOT send a copy of this form.

3/1/95



PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/019,596A

DATE: 02/03/2003 (-)
TIME: 10:31:18

Input Set : A:\PTO.DC.txt .

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3 <110> APPLICANT: University of Zurich
      5 <120> TITLE OF INVENTION: Hetero-associating coiled coil peptides
      7 <130> FILE REFERENCE: D 2398 PCT
      9 <140> CURRENT APPLICATION NUMBER: US/10/019,596A
    10 <141> CURRENT FILING DATE: 2002-07-11
    .12 <160> NUMBER OF SEQ ID NOS: 36
     14 <170> SOFTWARE: PatentIn version 3.0
     16 <210> SEQ ID NO: 1
     17 <211> LENGTH: 32
     18 <212> TYPE: PRT
     19 <213> ORGANISM: Artificial Sequence
     21 <220> FEATURE:
    22 <221> NAME/KEY: PEPTIDE
     23 <222> LOCATION: (1)..(32)
    24 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
     25 .
              construct
     27 <220> FEATURE:
W--> 28 <221> NAME/KEY: PEPTDIE
     29 <222> LOCATION: (1)..(32)
     30 <223> OTHER INFORMATION: Xaa at positions 5, 7, 12, 14, 19, 21, 26 and
             28 represents a mixture of glu, lys, gln or arg
     33 <220> FEATURE:
W--> 34 <221> NAME/KEY: PEPTDIE
     35 <222> LOCATION: (1)..(32)
     36 <223> OTHER INFORMATION: Xaa at position 15 represents a mixture of asn or val
     38 <400> SEQUENCE: 1
W--> 39 Val Ala Gln Leu Xaa Glu Xaa Val Lys Thr Leu Xaa Ala Xaa Xaa Tyr
     40 1
              . 5
                                            10
W--> 42 Glu Leu Xaa Ser Xaa Val Gln Arg Leu Xaa Glu Xaa Val Ala Gln Leu
                                        25
     47 <210> SEQ ID NO: 2
    48 <211> LENGTH: 32
     49 <212> TYPE: PRT.
    50 <213> ORGANISM: Artificial Sequence
    52 <220> FEATURE:
    53 <221> NAME/KEY: PEPTIDE .
    54 <222> LOCATION: (1)..(32)
   55 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic construct
    57 <220> FEATURE:
    58 <221> NAME/KEY: PEPTIDE
   · 59 <222> LOCATION: (1)..(32)
    60 <223> OTHER INFORMATION: Xaa at positions 5, 7, 12, 14, 19, 21, 26 and
             28 represents a mixture of glu, lys, gln or arg
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Input Set : A:\PTO.DC.txt

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64 <220> FEATURE:
    65 <221> NAME/KEY: PEPTIDE
     66 <222> LOCATION: (1)..(32)
    67 <223> OTHER INFORMATION: Xaa at position 15 represents a mixture of asn or val
    69 <400> SEQUENCE: 2
W--> 70 Val Asp Glu Leu Xaa Ala Xaa Val Asp Gln Leu Xaa Asp Xaa Xaa Tyr .
W--> 73 Ala Leu Xaa Thr Xaa Val Ala Gln Leu Xaa Lys Xaa Val Glu Lys Leu
            20
    78 <210> SEQ ID NO: 3
    79 <211> LENGTH: 32
    80 <212> TYPE: PRT ·
    81 <213> ORGANISM: artificial sequence
    83 <220> FEATURE:
    84 <221> NAME/KEY: PEPTIDE
    85 <222> LOCATION: (1)..(32)
    86 <223> OTHER INFORMATION: hetero-associating (poly)peptide
    88 <400> SEQUENCE: 3
    90 Val Ala Gln Leu Glu Glu Lys Val Lys Thr Leu Arg Ala Gln Asn Tyr
           5 10
    93 Glu Leu Lys Ser Arg Val Gln Arg Leu Arg Glu Gln Val Ala Gln Leu
                                      25
    97 <210> SEQ ID NO: 4
    98 <211> LENGTH: 32
    99 <212> TYPE: PRT
    100 <213> ORGANISM: artificial sequence
    102 <220> FEATURE:
    103 <221> NAME/KEY: PEPTIDE
    104 <222> LOCATION: (1)..(32)
    105 <223> OTHER INFORMATION: hetero-associating (poly) peptide
    107 <400> SEQUENCE: 4
    109 Val Ala Gln Leu Arg Glu Arg Val Lys Thr Leu Arg Ala Gln Asn Tyr
    112 Glu Leu Glu Ser Glu Val Gln Arg Leu Arg Glu Gln Val Ala Gln Leu
         20
                                       25
    116 <210> SEQ ID NO: 5
    117 <211> LENGTH: 32
    118 <212> TYPE: PRT
    119 <213> ORGANISM: artificial sequence
    121 <220> FEATURE:
    122 <221> NAME/KEY: PEPTIDE
    124 <222> LOCATION: (1)..(32)
    125 <223> OTHER INFORMATION: hetero-associating (poly)peptide
    127 <400> SEQUENCE: 5
    129 Val Ala Gln Leu Gln Glu Lys Val Lys Thr Leu Arg Ala Arg Asn Tyr
                                          10
   132 Glu Leu Lys Ser Glu Val Gln Arg Leu Glu Glu Lys Val Ala Gln Leu
    136 <210> SEQ ID NO: 6
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Input Set : A:\PTO.DC.txt

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137 <211> LENGTH: 32
  138 <212> TYPE: PRT
  139 <213> ORGANISM: artificial sequence
  141 <220> FEATURE:
  142 <221> NAME/KEY: PEPTIDE
  143 <222> LOCATION: (1)..(32)
  144 <223> OTHER INFORMATION: hetero-associating (poly)peptide
  146 <400> SEQUENCE: 6
  148 Val Ala Gln Leu Glu Glu Gln Val Lys Thr Leu Gln Ala Arg Asn Tyr
  149 1 .5
                              . 10
  151 Glu Leu Lys Ser Lys Val Gln Arg Leu Lys Glu Lys Val Ala Gln Leu
     20
  152
  155 <210> SEO ID NO: 7
  156 <211> LENGTH: 32.
  157 <212> TYPE: PRT
  158 <213> ORGANISM: artificial sequence
  160 <220> FEATURE:
  161 <221> NAME/KEY: PEPTIDE
  162 <222> LOCATION: (1)..(32)
 163 <223> OTHER INFORMATION: hetero-associating (poly)peptide
  165 <400> SEQUENCE: 7
  167 Val Ala Gln Leu Glu Glu Arg Val Lys Thr Leu Arg Ala Gln Asn Tyr
  168 1 5 10 15
 170 Glu Leu Lys Ser Lys Val Gln Arg Leu Glu Glu Gln Val Ala Gln Leu
 171 20 25
 174 <210> SEQ ID NO: 8
 175 <211> LENGTH: 32
 176 <212> TYPE: PRT
 177 <213> ORGANISM: artificial sequence
 179 <220> FEATURE:
 180 <221> NAME/KEY: PEPTIDE
 181 <222> LOCATION: (1)..(32) -
 182 <223> OTHER INFORMATION: hetero-associating (poly)peptide
 184 <400> SEQUENCE: 8
 186 Val Ala Gln Leu Glu Glu Gln Val Lys Thr Leu Glu Ala Glu Asn Tyr
 187 1 5 10
. 189 Glu Leu Lys Ser Lys Val Gln Arg Leu Arg Glu Arg Val Ala Gln Leu
 19Ó
     20
 193 <210> SEQ ID NO: 9
 194 <211> LENGTH: 32
 195 <212> TYPE: PRT
 196 <213> ORGANISM: artificial sequence
 198 <220> FEATURE:
 199 <221> NAME/KEY: PEPTIDE
 200 <222> LOCATION: (1)..(32)
 201 <223> OTHER INFORMATION: hetero-associating (poly) peptide
203 <400> SEQUENCE: 9
 205 Val Ala Gln Leu Gln Glu Gln Val Lys Thr Leu Glu Ala Gln Asn Tyr
 206 1
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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF4\02032003\J019596A.raw

208 Ģlu Leu Glu Ser Glu Val Gln Arg Leu Lys Glu Gln Val Ala Gln Leu 25 212 <210> SEQ ID NO: 10 213 <211> LENGTH: 32 214 <212> TYPE: PRT 215 <213> ORGANISM: artificial sequence 217 <220> FEATURE: 218 <221> NAME/KEY: PEPTIDE 219 <222> LOCATION: (1).:(32) 220 <223> OTHER INFORMATION: hetero-associating (poly)peptide 222 <400> SEQUENCE: 10 224 Val Ala Gln Leu Glu Glu Arg Val Lys Thr Leu Lys Ala Glu Asn Tyr 10 227 Glu Leu Glu Ser Glu Val Gln Arg Leu Lys Glu Arg Val Ala Gln Leu 20 231 <210> SEQ ID NO: 11 232 <211> LENGTH: 32 233 <212> TYPE: PRT 234 <213> ORGANISM: artificial sequence 236 <220> FEATURE: 237 <221> NAME/KEY: PEPTIDE 238 <222> LOCATION: (1)..(32) 239 <223> OTHER INFORMATION: hetero-associating (poly)peptide 241 <400> SEQUENCE: 11 244 Val Ala Gln Leu Glu Glu Lys Val Lys Thr Leu Lys Ala Lys Asn Tyr 245 1 · 10 247 Glu Leu Lys Ser Lys Val Gln Arg Leu Lys Glu Lys Val Ala Gln Leu 248 . 20 . 25 . 30 251 <210> SEQ ID NO: 12 252 <211> LENGTH: 32 253 <212> TYPE: PRT 254 <213> ORGANISM: artificial sequence 256 <220> FEATURE: 257 <221> NAME/KEY: PEPTIDE 258 <222> LOCATION: (1)..(32) 259 <223> OTHER INFORMATION: hetero-associating (poly)peptide 261 <400> SEQUENCE: 12 263 Val Ala Gln Leu Gln Glu Glu Val Lys Thr Leu Gln Ala Glu Asn Tyr 5 10 266 Glu Leu Arg Ser Glu Val Gln Arg Leu Glu Glu Glu Val Ala Gln Leu 270 <210> SEQ ID NO: 13 271 <211> LENGTH: 32 272 <212> TYPE: PRT 273 <213> ORGANISM: artificial sequence 275 <220> FEATURE: 276 <221> NAME/KEY: PEPTIDE 277 <222> LOCATION: (1)..(32) 278 <223> OTHER INFORMATION: hetero-associating (poly)peptide

Input Set : A:\PTO.DC.txt

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280 <400> SEQUENCE: 13
  282 Val Ala Gln Leu Arg Glu Arg Val Lys Thr Leu Arg Ala Arg Asn Tyr
  283 1 5 10 15
  285 Glu Leu Gln Ser Lys Val Gln Arg Leu Lys Glu Arg Val Ala Gln Leu
 286 20 25
 289 <210> SEQ ID NO: 14
 290 <211> LENGTH: 32
 291 <212> TYPE: PRT
 292 <213> ORGANISM: artificial sequence
 294 <220> FEATURE:
 295 <221> NAME/KEY: PEPTIDE
 296 <222> LOCATION: (1)..(32)
 297 <223> OTHER INFORMATION: hetero-associating (poly)peptide
 299 <400> SEQUENCE: 14
301 Val Asp Glu Leu Gln Ala Glu Val Asp Gln Leu Gln Asp Glu Asn Tyr
 302 1 5 10 ... 15
 304 Ala Leu Lys Thr Lys Val Ala Gln Leu Arg Lys Lys Val Glu Lys Leu
 305 20 25
 308 <210> SEQ ID NO: 15
 309 <211> LENGTH: 32
 310 <212> TYPE: PRT
 311 <213> ORGANISM: artificial sequence
 313 <220> FEATURE:
 314 <221> NAME/KEY: PEPTIDE
 315 <222> LOCATION: (1)..(32)
 316 <223> OTHER INFORMATION: hetero-associating (poly)peptide
 318 <400> SEQUENCE: 15
 320 Val Asp Glu Leu Lys Ala Glu Val Asp Gln Leu Gln Asp Gln Asn Tyr
 321 1 5 10 15
323 Ala Leu Arg Thr Lys Val Ala Gln Leu Arg Lys Glu Val Glu Lys Leu
 324 20 25 30
 327 <210> SEQ ID NO: 16
 328 <211> LENGTH: 32
 329 <212> TYPE: PRT
 330 <213> ORGANISM: artificial sequence
 332 <220> FEATURE:
 333 <221> NAME/KEY: PEPTIDE
 334 <222> LOCATION: (1)..(32)
 335 <223> OTHER INFORMATION: hetero-associating (poly)peptide
337 <400> SEQUENCE: 16
 339 Val Asp Glu Leu Glu Ala Glu Val Asp Gln Leu Lys Asp Gln Asn Tyr
 340 1 5 10
 342 Ala Leu Lys Thr Lys Val Ala Gln Leu Gln Lys Gln Val Glu Lys Leu
 343 20
 346 <210> SEQ ID NO: 17
 347 <211> LENGTH: 32
· 348 <212> TYPE: PRT
 349 <213> ORGANISM: artificial sequence
 351 <220> FEATURE:
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/019,596A

DATE: 02/03/2003 TIME: 10:31:20

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF4\02032003\J019596A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 5,7,12,14,15,19,21,26,28

Seq#:2; Xaa Pos. 5,7,12,14,15,19,21,26,28

Seq#:25; N Pos. 17,18,19,23,24,25,38,39,40,44,45,46,47,48,49,59,60,61,65,66

Seq#:25; N Pos. 67,79,80,81,85,86,87

Seq#:26; Xaa Pos. 5,7,12,14,15,19,21,26,28

Seq#:27; N Pos. 17,18,19,23,24,25,38,39,40,44,45,46,47,48,49,59,60,61,65,66

Seq#:27; N Pos. 67,79,80,81,85,86,87

Seq#:28; Xaa Pos. 5,7,12,14,15,19,21,26,28

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/019,596A

DATE: 02/03/2003 TIME: 10:31:20

Input Set : A:\PTO.DC.txt

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L:28 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:34 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:16
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:60
L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:570 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
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PCT10

Does Not Comply Corrected Dickette Needed

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/019,596A

DATE: 01/23/2003 TIME: 13:47:15

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF4\01232003\J019596A.raw

- 3 <110> APPLICANT: University of Zurich
- 5 <120> TITLE OF INVENTION: Hetero-associating coiled coil peptides
- 7 <130> FILE REFERENCE: D 2398 PCT
- 9 <140> CURRENT APPLICATION NUMBER: US/10/019,596A
- 10 <141> CURRENT FILING DATE: 2002-07-11
- 12 <160> NUMBER OF SEQ ID NOS: 36
- 14 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

- 720 <210> SEQ ID NO: 36 721 <211> LENGTH: 37
- 722 <212> TYPE: PRT
- 724 <213> ORGANISM: artificial sequence
- 726 <220> FEATURE:
- 727 <221> NAME/KEY: PEPTIDE
- 728 <222> LOCATION: (1)..(37)
- 729 <223> OTHER INFORMATION: N-acetylated and C-amidated synthetic peptide

10

٠. .

- 731 <400> SEQUENCE: 36
- 733 Ser Thr Ser Val Asp Glu Leu Gln Ala Glu Val Asp Gln Leu Gln Asp
- 5 ,
- 736 Glu Asn Tyr Ala Leu Lys Thr Lys Val Ala Gln Leu Arg Lys Val
- 737 20 25 30
- 739 Glu Lys Leu Ser Glu
- 740 35

E--> 755 (8) delete

VERIFICATION SUMMARY

DATE: 01/23/2003 PATENT APPLICATION: US/10/019,596A TIME: 13:47:16

Input Set : A:\pto.vsk.txt

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L:28 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:34 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:16
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:16
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:60 L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:16
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:570 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:60
L:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:16
L:755 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:36
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